

Amendments to the Specification:

Please replace the title on page 1 of the application with the following amended title:

**B7-H3 AND B7-H4, NOVEL A T CELL IMMUNOREGULATORY MOLECULES
MOLECULE**

Please replace the paragraph beginning on page 14, line 2, of the specification with the following amended paragraph:

The determination of percent identity between two sequences is accomplished using the mathematical algorithm of Karlin and Altschul, *Proc. Natl. Acad. Sci. USA* 90, 5873-5877, 1993. Such an algorithm is incorporated into the BLASTN and BLASTP programs of Altschul et al. (1990) *J. Mol. Biol.* 215, 403-410. BLAST nucleotide searches are performed with the BLASTN program, score = 100, wordlength = 12 to obtain nucleotide sequences homologous to B7-H3- or B7-H4-encoding nucleic acids. BLAST protein searches are performed with the BLASTP program, score = 50, wordlength = 3 to obtain amino acid sequences homologous to B7-H3 or B7-H4. To obtain gapped alignments for comparative purposes, Gapped BLAST is utilized as described in Altschul et al. (1997) *Nucleic Acids Res.* 25, 3389-3402. When utilizing BLAST and Gapped BLAST programs, the default parameters of the respective programs (e.g., XBLAST and NBLAST) are used (See <http://www.ncbi.nlm.nih.gov>).

Please replace the abstract at page 42 with the following amended abstract:

The invention provides ~~novel B7-H3 and B7-H4 polypeptides useful for co-stimulating T cells~~, isolated nucleic acid molecules ~~encoding them~~, vectors containing the nucleic acid molecules, and cells containing the vectors. Also included are methods of making ~~and using these B7-H4~~ co-stimulatory polypeptides.